

# The compartment bag test (CBT) for enumerating fecal indicator bacteria: basis for design and interpretation of results

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## Abstract

For the past several years, the compartment bag test (CBT) has been employed in water quality monitoring and public health protection around the world. To date, however, the statistical basis for the design and recommended procedures for enumerating fecal indicator bacteria (FIB) concentrations from CBT results have not been formally documented. Here, we provide that documentation following protocols for communicating the evolution of similar water quality testing procedures. We begin with an overview of the statistical theory behind the CBT, followed by a description of how that theory was applied to determine an optimal CBT design. We then provide recommendations for interpreting CBT results, including procedures for estimating quantiles of the FIB concentration probability distribution, and the confidence of compliance with recognized water quality guidelines. We synthesize these values in custom user-oriented ‘look-up’ tables similar to those developed for other FIB water quality testing methods. Modified versions of our tables are currently distributed commercially as part of the CBT testing kit.

### *Key words:*

Compartment bag test; water quality; drinking water; human health; statistical methods

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# 1 Introduction

2 Ensuring readily-available high quality drinking water is fundamental to hu-  
3 man health and has important connections to socioeconomic status, commer-  
4 cial and industrial growth, and overall quality of life (Mekonnen and Hoek-  
5 stra, 2016). The challenge of providing that ensurance is met in different ways  
6 around the world; in some communities, drinking water supplies are assumed  
7 protected if they are adequately separated from wastewater and other sources  
8 of contamination (George, 2008). In others, routine water quality testing is  
9 used to ensure compliance with recognized standards (Gleick, 1998; Novotny,  
10 2003). Testing kits that support these assessments often require a skilled tech-  
11 nician to collect, analyze, and interpret results, as well as microbiological lab-  
12 oratory facilities. In regions of the world without these resources and where  
13 the time from water withdrawal (from its source) to consumption is short,  
14 alternative testing procedures are needed.

15 To address this gap in global water quality protection, researchers at the Uni-  
16 versity of North Carolina Chapel Hill and Duke University developed a simple  
17 kit for enumerating FIB concentrations that is portable, relatively inexpen-  
18 sive, and provides easy-to-interpret results (Stauber et al., 2014). This kit,  
19 commonly referred to as the compartment bag test (or CBT), is currently  
20 manufactured and distributed by Aquagenx, LLC and has been tested and  
21 used in communities around the world (Murcott et al., 2015; Weiss et al.,  
22 2016). To date, however, the statistical basis for the design and recommended  
23 interpretation of results from the CBT have not been formally documented.

24 Here, following documentation for the development of similar water quality  
25 testing kits (McCrary, 1915; de Man, 1977; Tillett and Coleman, 1985; Haas,  
26 1989; McBride et al., 2003), we begin with an overview of the statistical the-

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27 ory behind the CBT, followed by examples of how that theory was applied  
28 to determine an optimal CBT design. We then provide recommendations for  
29 interpreting CBT results, including procedures for estimating quantiles of the  
30 FIB concentration probability distribution, as well as procedures for calcu-  
31 lating the confidence of compliance with World Health Organization (WHO)  
32 drinking water quality guidelines (McBride and Ellis, 2001; Borsuk et al.,  
33 2002; World Health Organization, 2004). We synthesize these values in cus-  
34 tom user-oriented ‘look-up’ tables similar to those developed for other FIB  
35 testing kits (de Man, 1977). Finally, we explore the sensitivity of CBT results  
36 to departures from assumptions in the underlying statistical models, and from  
37 recommended protocols for sample collection and handling.

## 38 2 Experimental

### 39 2.1 Statistical basis for interpreting CBT results

40 The CBT is a manufactured clear plastic multi-compartment bag into which  
41 100 ml of a water sample is distributed (Stauber et al., 2014). Each com-  
42 partment contains a growth substrate designed to detect groups of FIB (such  
43 as hydrogen sulphide producers), or specific bacteria such as *Escherichia coli*  
44 (EC), by turning a distinctive color (e.g. blue-green) indicating growth of “tar-  
45 get” (e.g. FIB or EC) bacteria during an incubation period. The CBT will  
46 yield a pattern of ‘positive’ and ‘negative’ compartments from which a user  
47 can infer the FIB concentration of the original sample following the common  
48 assumption (Greenwood and Yule, 1917; Cochran, 1950; Woodward, 1957;  
49 El-Shaarawi et al., 1981; Hurley and Roscoe, 1983; de Man, 1983; Haas and  
50 Heller, 1988; Woomer et al., 1990; Briones and Reichardt, 1999) that, for a  
51 given sample, the number of target bacteria ( $y_i$ ) in compartment  $i$  ( $i \in [1, m]$   
52 and  $m$  is the total number of compartments) with volume  $v_i$  (assuming a  
53 well-mixed sample) is well-represented by a Poisson probability distribution

54  $y_i \sim \text{Po}(\lambda_i = cv_i/100)$  with FIB concentration  $c$  (in organisms per 100 ml),  
 55 and mean and variance  $\lambda_i$ . The probability of a positive compartment of vol-  
 56 ume  $v_i$  is  $1 - \exp(-cv_i/100)$ . The joint probability of any pattern of positive and  
 57 negative compartments  $\vec{x}$  (where the over-arrow superscript denotes a row  
 58 vector,  $x_i \in [0,1]$  and  $x=1$  indicates a positive compartment) is then expressed  
 59 as the product of a series of  $m$  independent Bernoulli trials:

$$f(\vec{x} | \vec{v}, c) \propto \prod_{i=1}^m \left(1 - e^{-cv_i/100}\right)^{x_i} \left(e^{-cv_i/100}\right)^{1-x_i} \quad (1)$$

60 Conventional interpretations of presence/absence test kits for FIB often focus  
 61 on a deterministic solution to  $c$  from equation 1. This value is commonly  
 62 referred to as the “most probable number” (or MPN) and can be calculated  
 63 as (Hurley and Roscoe, 1983; McBride, 2005; Gronewold and Wolpert, 2008):

$$\text{MPN} = \underset{c}{\text{argmax}} \left[ \prod_{i=1}^m \left(1 - e^{-cv_i/100}\right)^{x_i} \left(e^{-cv_i/100}\right)^{1-x_i} \right] \quad (2)$$

64 We implement this formulation using the `uniroot` function in the R statistical  
 65 software package (R core team, 2014). Corresponding code is included in the  
 66 Supplementary Information.

67 Multiple methods have been developed for expressing uncertainty in the MPN,  
 68 however most do not explicitly acknowledge that the probability distribution  
 69 of the MPN for a given pattern of positive and negative compartments is  
 70 typically discrete and multi-modal, while the probability distribution of the  
 71 FIB concentration is almost always unimodal and continuous (Klee, 1993;  
 72 Gronewold and Wolpert, 2008). Therefore, in addition to reporting conven-  
 73 tional MPN values, we propose two interpretations of CBT results that allow  
 74 for a more robust understanding of the uncertainty in the FIB concentration  
 75 and how that uncertainty affects the confidence of compliance with water qual-  
 76 ity guidelines (McBride and Ellis, 2001; Gronewold and Borsuk, 2009, 2010).

77 The first is based on calculating quantiles of the likelihood function of the FIB  
78 concentration (equation 1, written as a function of  $c$  for given  $\vec{x}$  and  $\vec{v}$ ), as  
79 well as the probability that the FIB concentration exceeds 1, 10, 100, or 1000  
80 organisms per 100 ml.

81 The second interpretation is based on a Bayesian analysis of CBT results  
82 (Bernardo and Ramon, 1998; Press, 2003; Bolstad, 2004) where the posterior  
83 probability distribution of the FIB concentration  $c$  is proportional to the prod-  
84 uct of the likelihood function (equation 1) and prior probability distribution  
85  $\pi(c)$ :

$$f(c | \vec{x}, \vec{v}) \propto \pi(c)f(\vec{x} | \vec{v}, c) \quad (3)$$

86 One advantage of this approach is that it allows for expression of *a priori*  
87 assumptions about the potential range of the FIB concentration in a water  
88 sample. Methods based on the likelihood function alone, in contrast, implicitly  
89 assume *a priori* that FIB concentrations ranging from 0 to  $\infty$  are equally  
90 likely; an assumption analogous to a belief that gross contamination is just  
91 as likely as a FIB concentration within a few orders of magnitude of (or even  
92 well below) WHO water quality guidelines. This *a priori* belief is just one of  
93 many a CBT user might have about water quality at a particular sampling  
94 location (Press, 2003). Here, we present calculations based on a lognormal  
95 prior  $\pi(c) = \text{LN}(\mu = 0, \sigma^2 = 100)$ , with log-concentration mean  $\mu$  and variance  
96  $\sigma^2$ , intended to represent an *a priori* belief that the FIB concentration is most  
97 likely low, but that extreme FIB concentrations are possible. We view further  
98 investigation of impacts of alternative priors on CBT results as an important  
99 area for future research.

100 It is informative to note that previous studies have explored alternative prob-  
101 ability models for interpreting multiple-compartment water quality analysis  
102 results, including the negative binomial model and variations of the Poisson

103 model that account for thinning and dispersion (Christian and Pipes, 1983;  
104 El-Shaarawi et al., 1981; Messner and Wolpert, 2002; Crainiceanu et al., 2003).  
105 Recent research, however (see Gronewold et al., 2008; Wu et al., 2014), indi-  
106 cates that only extreme and persistent violations of the Poisson probability  
107 model would justify application of an alternative probability model.

108 Finally, following equation 1, we calculate the relative likelihood of each possi-  
109 ble combination of positive and negative compartments. Results of this calcu-  
110 lation provide an indication of CBT outcomes that are most likely, and those  
111 that (because they are extremely unlikely) might indicate contamination or  
112 thinning of individual compartments and would therefore warrant additional  
113 testing and verification.

## 114 *2.2 Design criteria*

115 The number and volume of compartments of the CBT is based on consid-  
116 eration of a range of criteria including ease of manufacturing, minimization  
117 of potential user error (such as unintentionally distributing more or less wa-  
118 ter into each CBT compartment than intended), and results that are readily  
119 translatable into health risk-based metrics. More specifically, the ideal CBT  
120 design yields a pattern of positive and negative compartments that are easy  
121 to translate into FIB concentrations with uncertainty bounds relevant to hu-  
122 man health risks. For most applications of the CBT, we expect these risks  
123 will be assessed using FIB concentration numeric limits prescribed in WHO  
124 water quality guidelines. We assess compliance with this criteria by inferring  
125 FIB concentrations associated with each possible result (i.e. each combina-  
126 tion of positive and negative compartments) of a particular CBT design, and  
127 then comparing these concentrations to established water quality criteria and  
128 standards.

129 To demonstrate our approach, we provide a comparison between two CBT

130 designs. The first (the design ultimately employed in practice) is a CBT with  
131 five compartments with volumes (in ml)  $\vec{v} = \{56, 30, 10, 3, 1\}$ . The second  
132 is a CBT with seven compartments with volumes  $\vec{v} = \{37, 32, 16, 8, 4, 2, 1\}$ .  
133 These design options evolved out of a qualitative consideration of the afore-  
134 mentioned criteria, as well as the constraints that the cumulative volume of  
135 all compartments equal 100ml, and that the compartment volumes span as  
136 broad a range as possible without multiple compartments of the same volume.

137 For each of the two test designs, we first calculated the full FIB concentration  
138 likelihood function for each possible CBT result, and then implemented our  
139 Bayesian interpretation by simulating samples from the posterior probability  
140 distribution of the FIB concentration (equation 3) for each possible CBT result  
141 using Markov chain Monte Carlo (MCMC) procedures in the software program  
142 WinBUGS (Lunn et al., 2000). We ran each MCMC chain until it reached  
143 convergence, indicated by a potential scale reduction factor  $\hat{R}$  (Gelman et al.,  
144 2004) close to 1.0. WinBUGS code used to simulate the posterior probability  
145 distribution for  $c$  for the  $\vec{v} = \{56, 30, 10, 3, 1\}$  CBT design is included in  
146 the Supplementary Information. From the likelihood functions and posterior  
147 probability distributions, we calculate a series of quantiles, as well as the  
148 likelihood (or posterior probability) that the FIB concentration exceeds 1, 10,  
149 100, or 1000 organisms per 100 ml.

### 150 *2.3 Sensitivity analysis*

151 To better understand the sensitivity of CBT results to potential variations  
152 in user handling (including violations of the assumptions in our statistical  
153 models), we repeat the simulation described in the previous section for the 5-  
154 compartment CBT using hypothetical compartment volumes (in ml) of  $\vec{v} =$   
155  $\{58.4, 30.5, 14.5, 2.5, 0.7\}$  and  $\vec{v} = \{32.3, 33.5, 23.3, 4.9, 3.4\}$ . These volume  
156 sequences were obtained from an informal (unpublished) study by one of the

157 authors at the University of North Carolina - Chapel Hill in which roughly  
158 twenty individuals with a range of CBT experience used the CBT, and the  
159 actual water sample volumes they distributed into each compartment were  
160 recorded. The two selected sequences represent, respectively, moderate and  
161 severe departures from the intended 5-compartment CBT design with com-  
162 partment volumes  $\vec{v} = \{56, 30, 10, 3, 1\}$ .

### 163 **3 Results and discussion**

164 Of the 32 potential combinations of positive and negative compartments for  
165 the 5-compartment CBT, we find that there are appreciable differences in the  
166 relative likelihood of each outcome (see Table S1 in Supplementary Informa-  
167 tion). Some results (particularly those for which the 56ml compartment is  
168 positive) are quite likely while others are highly improbable. This is an impor-  
169 tant distinction because highly unlikely CBT outcomes might indicate one or  
170 more potential problems with sample handling and analysis (including thin-  
171 ning or contamination of a particular compartment) and warrant additional  
172 investigation. To underscore this point, and to simplify our discussion of alter-  
173 native CBT interpretations, we hereafter focus on results from only the eight  
174 most likely outcomes of the CBT.

175 FIB concentration likelihood functions reflecting information content of indi-  
176 vidual CBT compartments (top five rows figure 1), and of each combination  
177 of positive and negative compartments for the eight most likely CBT results  
178 (bottom row figure 1), provide insight into origins of uncertainty in CBT-based  
179 water quality assessments (see also table 1). For example, a CBT result with  
180 a pattern of positive (1) and negative (0) compartments (with volumes 56,  
181 30, 10, 3, and 1ml) of  $\vec{x} = \{1, 1, 0, 1, 0\}$  has an MPN of 9.6 (organisms per  
182 100 ml) with moderate certainty in the FIB concentration. A CBT result for  
183 which the pattern of positive and negative compartments is  $\vec{x} = \{1, 1, 1, 1, 0\}$

184 has a higher MPN (48.3) and more uncertainty in the FIB concentration be-  
185 cause of the difference in the information content of the 10ml compartment.  
186 A positive 10ml compartment (by itself) indicates that the FIB concentration  
187 is almost certainly above roughly 40 organisms per 100 ml, while a negative  
188 10ml compartment indicates that the concentration is almost certainly be-  
189 low 40 organisms per 100ml. The contrast between the information in these  
190 two results underscores not only the relative value of keeping the CBT simple  
191 (by minimizing the number of compartments, for example) and easy to im-  
192 plement, but also the potential sensitivity of CBT outcomes to variations in  
193 sample handling.

194 A Bayesian interpretation of results from the 5-compartment CBT with  $\vec{v} =$   
195  $\{56, 30, 10, 3, 1\}$  (figure 2 and table 2) indicates how explicit quantification  
196 of *a priori* beliefs about the FIB concentration in a sample can propagate  
197 into different perceptions of human health risk (figure 2) when compared to  
198 interpretations based on the likelihood function alone, particularly for CBT  
199 outcomes with an intrinsically broad likelihood function (e.g. a result of  $\vec{x} =$   
200  $\{1, 1, 1, 1, 0\}$ ). In areas where there is a long history of high quality drinking  
201 water, for example, a prior probability distribution reflecting a strong belief  
202 in a relatively low FIB concentration may be helpful in guiding water use  
203 management decisions when there is insufficient information content in the  
204 likelihood function alone.

205 We also find that the 5-compartment CBT design (tables 1 and 2) provides a  
206 robust basis for distinguishing samples based on compliance with WHO water  
207 quality guidelines, particularly when compared to our alternative design with  
208 seven compartments (see table S2 in Supplementary Information). For nearly  
209 all of the most likely results of the 5-compartment CBT, we can make a rela-  
210 tively confident statement about the range of the sample FIB concentration,  
211 and about compliance with each numeric limit in the WHO guidelines. This  
212 statement may depend, as we have shown, on whether a likelihood or Bayesian

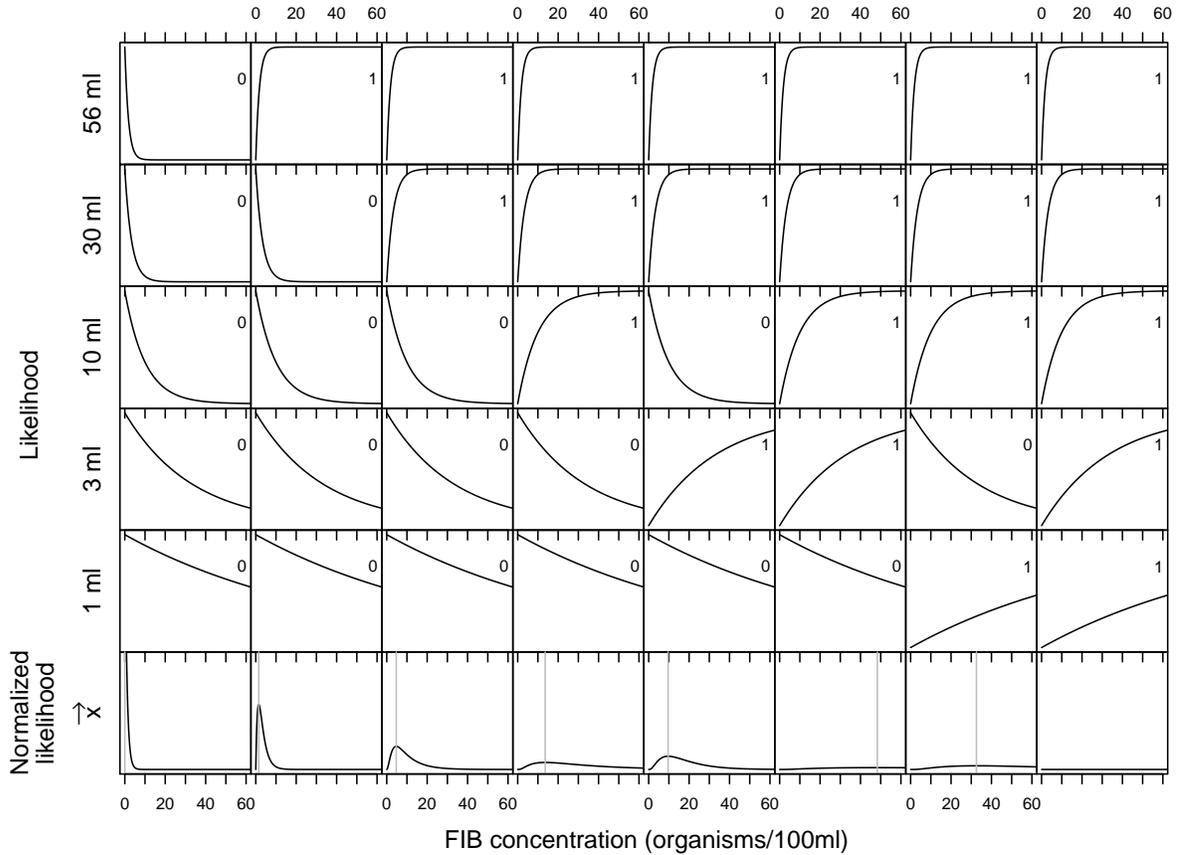


Fig. 1. Likelihood functions (first five rows) for individual positive ('1' in upper-right corner of panel) or negative ('0' in upper-right corner) compartments of the 5-compartment CBT, and normalized likelihood functions (bottom row) for the eight most likely outcomes of the CBT. Likelihood functions in the bottom row reflect the combined results of the positive and negative compartments from the five panels above (from the same column). The bottom left-most panel, for example, is the normalized likelihood function for the FIB concentration from a CBT result with all compartments negative. Vertical grey lines in the panels of the bottom row indicate the MPN (note that the MPN is undefined when all compartments are positive).

213 interpretation is used. In either case, a probabilistic interpretation enhances  
 214 information from conventional MPN values alone; water quality experts are  
 215 often comfortable with MPN values, but not with quantifying associated un-  
 216 certainties when the MPN is derived from a novel and unconventional testing  
 217 kit such as the CBT.

218 Our assessment of the potential impacts of user error (table 3 and Supple-  
 219 mentary Information) suggests that the 5-compartment CBT test is relatively  
 220 robust to both moderate and severe errors. More specifically, we find that

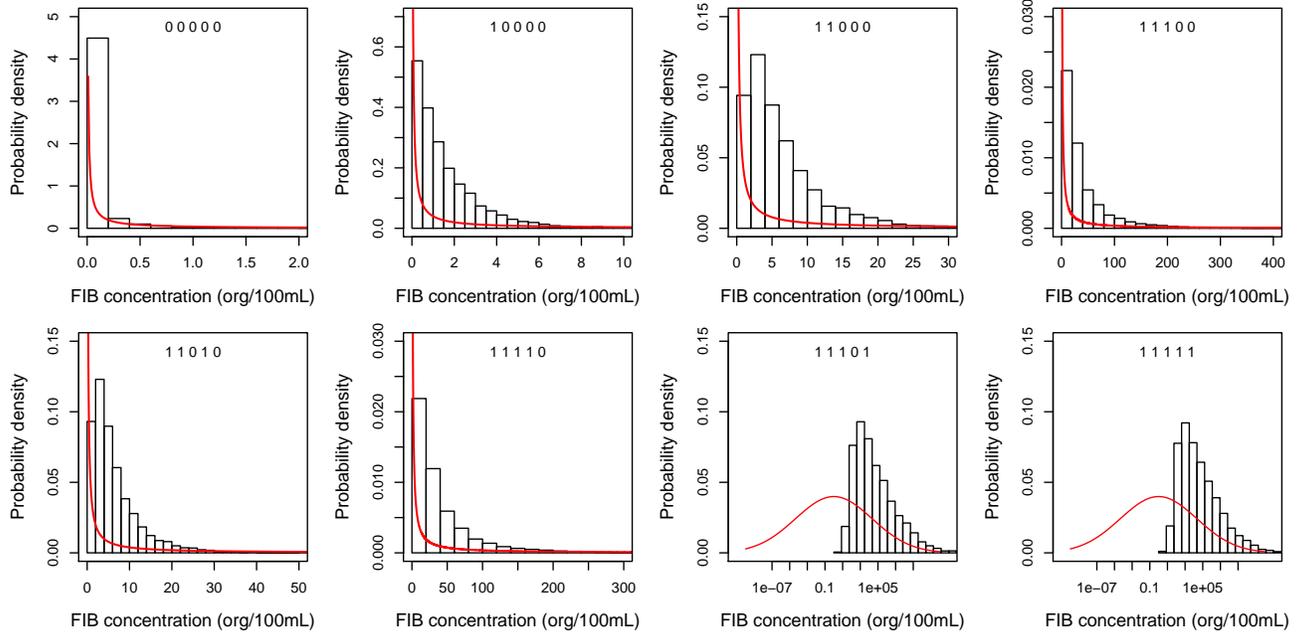


Fig. 2. Bayesian interpretation of CBT results including FIB concentration prior probability distribution (red lines) and histograms of simulated samples from the FIB concentration posterior probability distribution for the eight most likely results from the 5-compartment (volumes 56, 30, 10, 3, 1 ml) CBT. Values of 1 and 0 across the top of each panel correspond to each pattern of positive (1) and negative (0) compartments of volumes 56, 30, 10, 3, 1 ml, respectively. The x-axis of the two right-most panels in the bottom row is plotted on a logarithmic scale for clarity.

221 moderate handling errors would not have changed the perceived probability  
 222 of violating the WHO water quality guideline of 100 organisms per 100 ml  
 223 (a value indicating ‘very high risk’ water). Furthermore, we find that severe  
 224 errors, while leading to a slightly lower perceived probability of violating the  
 225 WHO water quality guideline of 100 organisms per 100 ml, would also have  
 226 been very unlikely to lead to a different perception of risk than what would  
 227 have been inferred had there been no error.

228 Finally, we acknowledge that users of the CBT have inquired about the un-  
 229 certainty in CBT results relative to uncertainties in more conventional water  
 230 quality testing tools, including (for example) membrane filtration (MF) tests  
 231 (Dufour and Cabelli, 1975; Dufour et al., 1981; El-Shaarawi et al., 1981). A  
 232 comparison between the 95% likelihood intervals from our analysis of the CBT  
 233 (table 1) and 95% likelihood intervals from MF tests with colony-forming unit

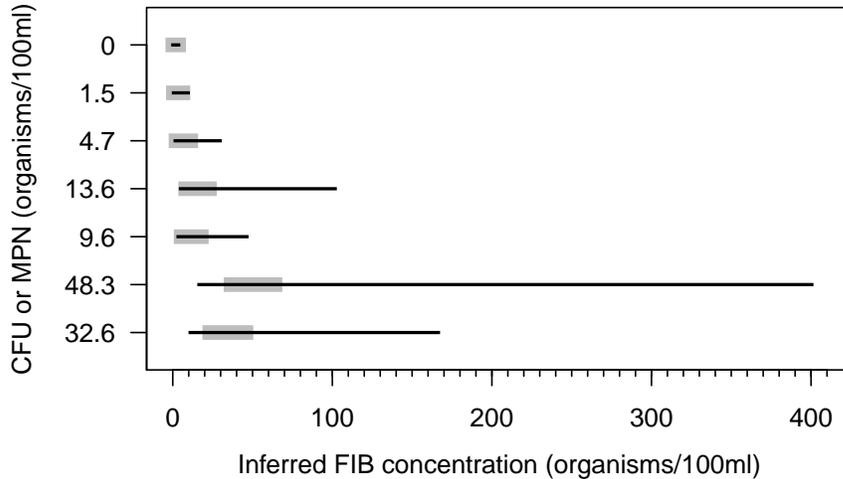


Fig. 3. FIB concentration 95% likelihood intervals based on seven of the eight most likely results of the 5-compartment CBT (intervals for a CBT result with all compartments positive are not shown because the likelihood function is continuously increasing and the MPN is undefined). Thin black segments represent likelihood intervals derived from the CBT. Thick grey segments represent likelihood intervals derived from conventional membrane filtration (MF) analyses with CFU values that correspond to MPN values from the CBT. The top-most pair of segments, for example, includes FIB concentration 95% likelihood intervals from (thin black segment) a CBT result with an MPN of 0, and (thick grey segment) an MF result with a CFU of 0.

234 (CFU) values matching MPN values from the CBT (Gronewold and Wolpert,  
 235 2008) indicates that (figure 3), for very low (i.e. less than 5 organisms per  
 236 100ml) FIB concentrations, the confidence intervals are quite similar and that  
 237 the differences are more extreme for FIB concentration close to and above  
 238 10 (organisms per 100ml). A Bayesian interpretation of CBT results (table  
 239 2) could affect the range of these intervals and might in fact be desirable  
 240 should water quality management officials (and other CBT users) find that  
 241 the likelihood-based intervals do not provide enough informative at higher  
 242 concentrations. We suggest investigation of impacts of alternative prior distri-  
 243 butions on inferred FIB concentration uncertainty and compliance with WHO  
 244 water quality guidelines as a high priority for future research.

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| 56mL | 30mL | 10mL | 3mL | 1mL | MPN  | $q_{2.5}$ | $q_{5.0}$ | $q_{25.0}$ | $q_{75.0}$ | $q_{95.0}$ | $q_{97.5}$ | Likelihood that $c >$ |      |      |      |
|------|------|------|-----|-----|------|-----------|-----------|------------|------------|------------|------------|-----------------------|------|------|------|
|      |      |      |     |     |      |           |           |            |            |            |            | 1                     | 10   | 100  | 1000 |
| 0    | 0    | 0    | 0   | 0   | 0.0  | <0.1      | <0.1      | 0.3        | 1.4        | 3.0        | 3.7        | 0.33                  | 0.00 | 0.00 | 0.00 |
| 1    | 0    | 0    | 0   | 0   | 1.5  | 0.4       | 0.5       | 1.5        | 4.4        | 8.1        | 9.7        | 0.85                  | 0.02 | 0.00 | 0.00 |
| 1    | 1    | 0    | 0   | 0   | 4.7  | 1.5       | 2.1       | 4.8        | 13.2       | 24.8       | 29.7       | 0.99                  | 0.39 | 0.00 | 0.00 |
| 1    | 1    | 1    | 0   | 0   | 13.6 | 4.8       | 6.4       | 15.1       | 44.1       | 84.4       | 101.8      | 1.00                  | 0.87 | 0.03 | 0.00 |
| 1    | 1    | 0    | 1   | 0   | 9.6  | 3.3       | 4.3       | 9.2        | 23.0       | 39.7       | 46.5       | 1.00                  | 0.71 | 0.00 | 0.00 |
| 1    | 1    | 1    | 1   | 0   | 48.3 | 16.4      | 22.4      | 55.5       | 170.0      | 331.0      | 400.5      | 1.00                  | 0.99 | 0.50 | 0.00 |
| 1    | 1    | 1    | 0   | 1   | 32.6 | 10.9      | 14.3      | 31.6       | 81.4       | 141.8      | 166.5      | 1.00                  | 0.98 | 0.16 | 0.00 |
| 1    | 1    | 1    | 1   | 1   | NA   | NA        | NA        | NA         | NA         | NA         | NA         | NA                    | NA   | NA   | NA   |

Table 1. Likelihood function-based interpretation of the eight highest likelihood combinations of positive (1) and negative (0) compartments for the recommended 5-compartment CBT design. Results include the MPN, quantiles of the FIB concentration normalized likelihood function, and the likelihood-based probability that  $c$  exceeds numeric water quality guidelines of 1, 10, 100, and 1000 (organisms per 100 ml). Note that when all compartments are positive, the FIB concentration likelihood function is continuously increasing and therefore the MPN and FIB concentration quantiles are undefined (Gronewold et al., 2010).

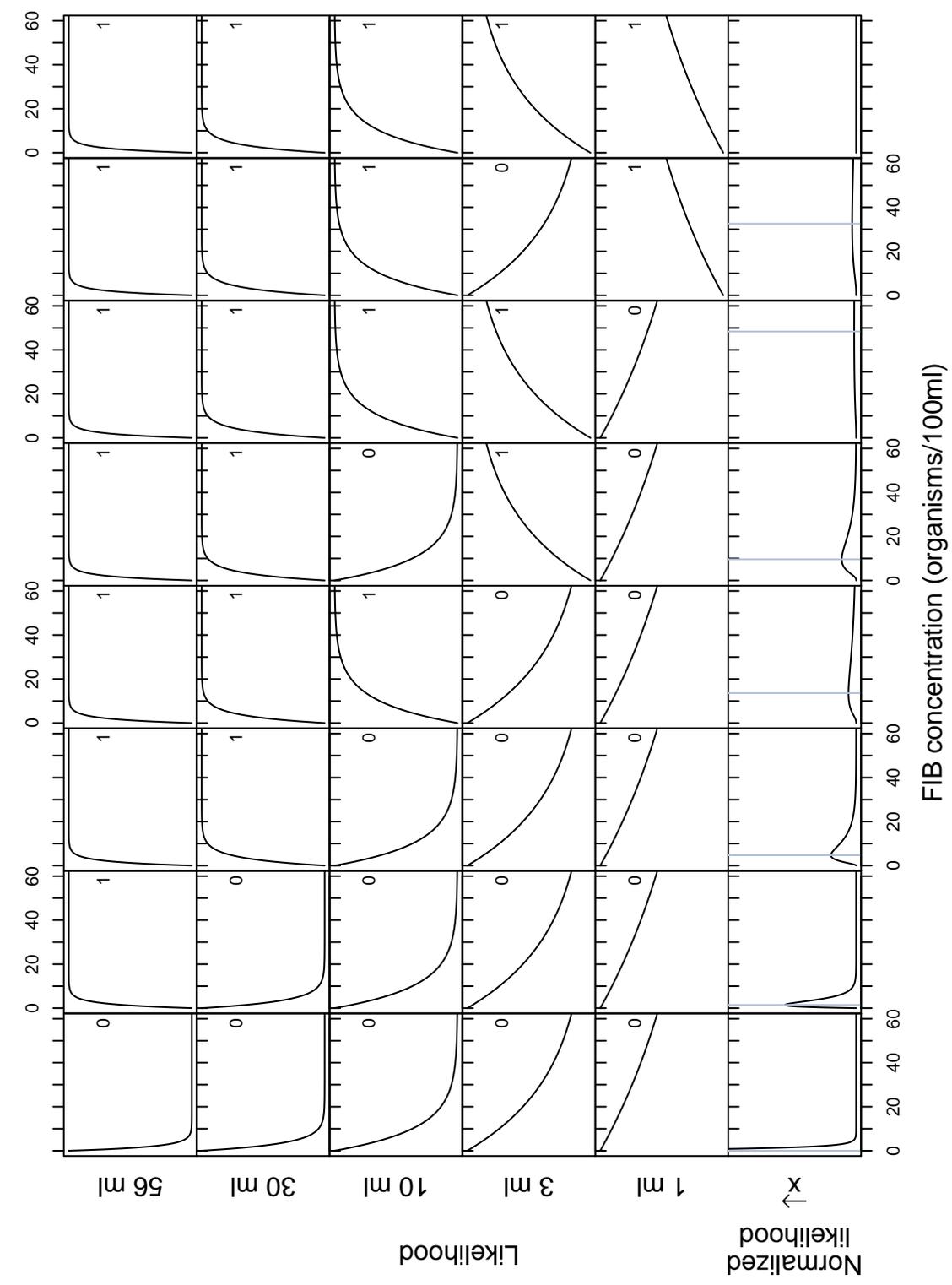
| 56mL | 30mL | 10mL | 3mL | 1mL | MPN  | $q_{2.5}$ | $q_{5.0}$ | $q_{25.0}$ | $q_{75.0}$        | $q_{95.0}$        | $q_{97.5}$           | Post. prob. that $c >$ |      |      |      |
|------|------|------|-----|-----|------|-----------|-----------|------------|-------------------|-------------------|----------------------|------------------------|------|------|------|
|      |      |      |     |     |      |           |           |            |                   |                   |                      | 1                      | 10   | 100  | 1000 |
| 0    | 0    | 0    | 0   | 0   | 0    | <0.1      | <0.1      | <0.1       | <0.1              | 0.5               | 0.8                  | 0.02                   | 0.00 | 0.00 | 0.00 |
| 1    | 0    | 0    | 0   | 0   | 1.5  | <0.1      | 0.1       | 0.4        | 2.2               | 4.9               | 6.2                  | 0.51                   | 0.00 | 0.00 | 0.00 |
| 1    | 1    | 0    | 0   | 0   | 4.7  | 0.6       | 0.9       | 2.5        | 8.3               | 18.1              | 22.0                 | 0.94                   | 0.18 | 0.00 | 0.00 |
| 1    | 1    | 1    | 0   | 0   | 13.6 | 2.8       | 3.8       | 11.1       | 49.7              | 142.5             | 189.3                | 1.00                   | 0.78 | 0.10 | 0.00 |
| 1    | 1    | 0    | 1   | 0   | 9.6  | 0.6       | 0.8       | 2.5        | 8.3               | 18.9              | 23.6                 | 0.94                   | 0.19 | 0.00 | 0.00 |
| 1    | 1    | 1    | 1   | 0   | 48.3 | 2.8       | 3.9       | 11.0       | 50.8              | 143.6             | 191.2                | 1.00                   | 0.78 | 0.10 | 0.00 |
| 1    | 1    | 1    | 0   | 1   | 32.6 | 37.4      | 70.7      | 734.7      | $8.5 \times 10^5$ | $1.2 \times 10^9$ | $1.9 \times 10^{10}$ | 1.00                   | 1.00 | 0.92 | 0.70 |
| 1    | 1    | 1    | 1   | 1   | NA   | 36.9      | 67.2      | 702.9      | $8.5 \times 10^5$ | $1.4 \times 10^9$ | $2.0 \times 10^{10}$ | 1.00                   | 1.00 | 0.93 | 0.73 |

Table 2. Bayesian interpretation of the eight highest likelihood combinations of positive (1) and negative (0) compartments for the recommended 5-compartment CBT design. Results include the MPN, quantiles of the FIB concentration posterior probability distribution, and the posterior probability that  $c$  exceeds numeric water quality guidelines of 1, 10, 100, and 1000 (organisms per 100 ml). Note that with a Bayesian interpretation, quantiles of the FIB concentration posterior probability distribution are defined when all compartments are positive, however the MPN is not defined (when all compartments are positive) because it is based on the likelihood function alone (Gronewold et al., 2010). A Bayesian interpretation of all possible combinations of positive and negative compartments is included in the Supplementary Information.

| Highest likelihood combinations of<br>pos. (1) and neg. (0) compartments |      |      |     |     | MPN  |      |      | $c_{95}$          |                   |                   | P( $c > 100$ ) |     |     |
|--|------|------|-----|-----|------|------|------|-------------------|-------------------|-------------------|----------------|-----|-----|
| 56mL   | 30mL | 10mL | 3mL | 1mL | *    | **   | ***  | *                 | **                | ***               | *              | **  | *** |
| 0  | 0    | 0    | 0   | 0   | 0.0  | 0.0  | 0.0  | 0.5               | 0.5               | 0.4               | 0.0            | 0.0 | 0.0 |
| 1  | 0    | 0    | 0   | 0   | 1.5  | 1.4  | 1.2  | 4.9               | 4.6               | 4.1               | 0.0            | 0.0 | 0.0 |
| 1  | 1    | 0    | 0   | 0   | 4.7  | 4.1  | 3.4  | 18.1              | 14.0              | 10.2              | 0.0            | 0.0 | 0.0 |
| 1  | 1    | 1    | 0   | 0   | 13.6 | 13.0 | 8.4  | 142.5             | 175.0             | 50.2              | 0.1            | 0.1 | 0.0 |
| 1  | 1    | 0    | 1   | 0   | 9.6  | 7.8  | 5.8  | 18.9              | 14.4              | 10.2              | 0.0            | 0.0 | 0.0 |
| 1  | 1    | 1    | 1   | 0   | 48.3 | 60.9 | 19.6 | 143.6             | 178.8             | 49.5              | 0.1            | 0.1 | 0.0 |
| 1  | 1    | 1    | 0   | 1   | 32.6 | 36.2 | 17.3 | $1.2 \times 10^9$ | $1.8 \times 10^9$ | $0.9 \times 10^9$ | 0.9            | 0.9 | 0.8 |
| 1  | 1    | 1    | 1   | 1   | NA   | NA   | NA   | $1.4 \times 10^9$ | $1.8 \times 10^9$ | $1.0 \times 10^9$ | 0.9            | 0.9 | 0.8 |

Table 3. Comparison between results when there is minimal (or no) user error (\*) and results with there is either moderate (\*\*) or severe (\*\*\*) user error. MPN values and 95<sup>th</sup> percentiles of the FIB concentration ( $c_{95}$ ) are in organisms per 100 ml. The final column indicates the posterior probability that the FIB concentration  $c$  exceeds the WHO numeric water quality standard of 100 organisms per 100 ml. Note that with a Bayesian interpretation, quantiles of the FIB concentration posterior probability distribution are defined when all compartments are positive, however the MPN is not defined (when all compartments are positive) because it is based on the likelihood function alone (Gronewold et al., 2010).

Figure 1  
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**Figure 2**  
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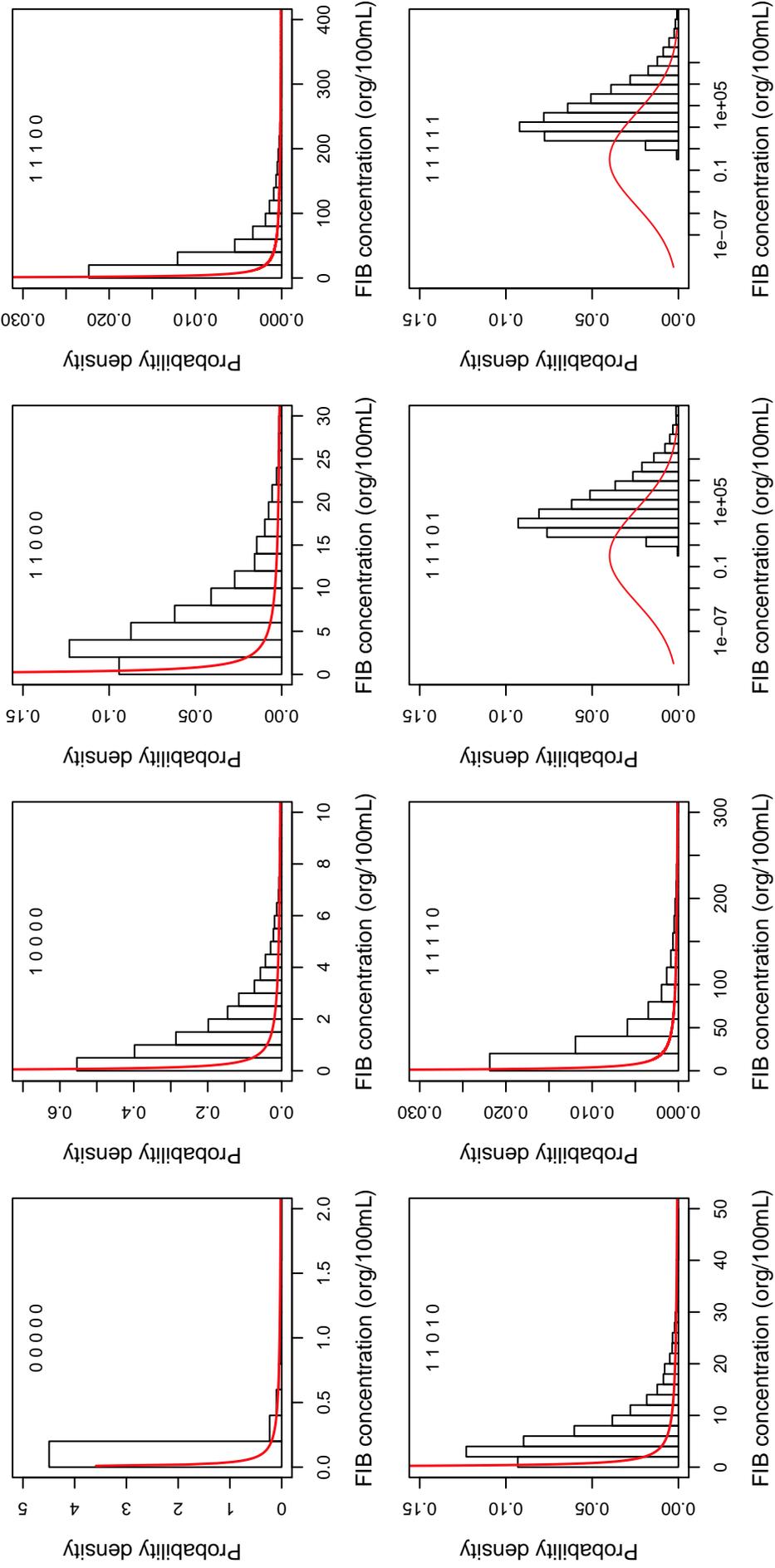
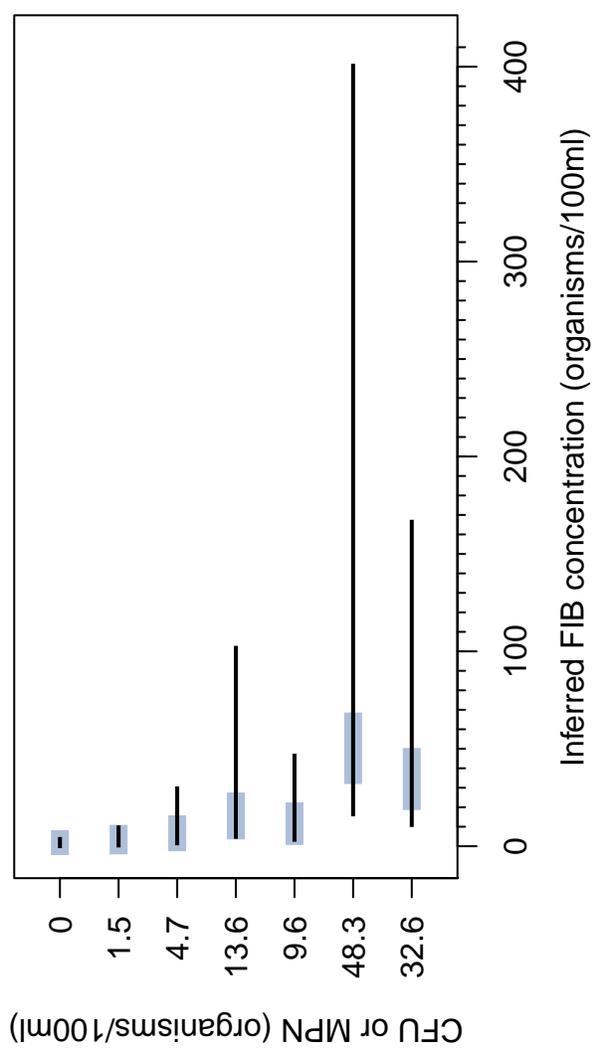


Figure 3  
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